

<210> 260  
<211> 22  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide probe

<400> 260  
caaagcctgc gcctggctctg tg 22

<210> 261  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide probe

<400> 261  
ttctggagcc cagagggtgc tgag 24

<210> 262  
<211> 45  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Synthetic oligonucleotide probe

<400> 262  
ggagctgccca cccattcaaa tggagcacga aggagagttc acctg 45

<210> 263  
<211> 2857  
<212> DNA  
<213> Homo sapiens

<400> 263  
tgaagagtaa tagttggaat caaaagagtc aacgcaatga actggtatatt 50  
actgctgcgt tttatgttgg gaattcctct cctatggcct tgtcttggag 100  
caacagaaaa ctctcaaaca aagaaagtca agcagccagt gcgatctcat 150  
ttgagagtga agcgtggctg ggtgtggaac caattttttg taccagagga 200  
aatgaatacg actagtcatc acatcggcca gctaagatct gatttagaca 250  
atggaaacaa ttctttccag tacaagcttt tgggagctgg agctggaagt 300  
acttttatca ttgatgaaag aacaggtgac atatatgccca tacagaagct 350  
tgatagagag gagcgatccc tctacatctt aagagcccag gtaatagaca 400  
tcgctactgg aagggtgtg gaacctgagt ctgagtttgt catcaaagtt 450

tcggatatca atgacaatga accaaaattc ctagatgaac cttatgaggc 500  
cattgtacca gagatgtctc cagaaggaac attagttatc caggtgacag 550  
caagtgatgc tgaogatccc tcaagtggta ataatgctcg tctcctctac 600  
agcttacttc aaggccagcc atatTTTTTt gttgaaccaa caacaggagt 650  
cataagaata tcttctaaaa tggatagaga actgcaagat gagtattggg 700  
taatcattca agccaaggac atgattggtc agccaggagc gttgtctgga 750  
acaacaagtg tattaattaa actttcagat gttaatgaca ataagcctat 800  
atttaaagaa agtttatacc gcttgactgt ctctgaatct gcacccactg 850  
ggacttctat aggaacaatc atggcatatg ataatgacat aggagagaat 900  
gcagaaatgg attacagcat tgaagaggat gattcgcaa catttgacat 950  
tattactaat catgaaactc aagaaggaat agttatatta aaaaagaaaag 1000  
tggattttga gcaccagaac cactacggta ttagagcaaa agttaaaaac 1050  
catcatgttc ctgagcagct catgaagtac cacactgagg cttccaccac 1100  
tttcattaag atccagggtg aagatgttga tgagcctcct cttttcctcc 1150  
ttccatatta tgtatttgaa gtttttgaag aaacccaca gggatcattt 1200  
gtaggcgtgg tgtctgccac agaccagac aataggaaat ctctatcag 1250  
gtattctatt actaggagca aagtgttcaa tatcaatgat aatggtaaca 1300  
tcaactacaag taactcactg gatcgtgaaa tcagtgcttg gtacaacct 1350  
agtattacag ccacagaaaa atacaatata gaacagatct cttcgatccc 1400  
actgtatgtg caagttctta acatcaatga tcatgctcct gagttctctc 1450  
aatactatga gacttatgtt tgtgaaaatg caggctctgg tcaggtaatt 1500  
cagactatca gtgcagtgga tagagatgaa tccatagaag agcaccattt 1550  
ttactttaat ctatctgtag aagacactaa caattcaagt tttaacaatca 1600  
tagataatca agataacaca gctgtcattt tgactaatag aactggtttt 1650  
aaccttcaag aagaacctgt cttctacatc tccatcttaa ttgccgacaa 1700  
tggaatcccg tcaattacaa gtacaaacac ccttaccatc catgtctgtg 1750  
actgtggtga cagtgggagc acacagacct gccagtacca ggagcttgtg 1800  
ctttccatgg gattcaagac agaagttatc attgctattc tcatttgcac 1850  
tatgatcata tttgggttta tttttttgac tttgggttta aaacaacgga 1900

gaaaacagat tctatttcct gagaaaagtg aagatttcag agagaatata 1950  
 ttccaatatg atgatgaagg ggggtggagaa gaagatacag aggcctttga 2000  
 tatagcagag ctgaggagta gtaccataat gcgggaacgc aagactcgga 2050  
 aaaccacaag cgctgagatc aggagcctat acaggcagtc tttgcaagtt 2100  
 ggccccgaca gtgccatatt caggaaattc attctggaaa agctcgaaga 2150  
 agctaatact gatccgtgtg cccctccttt tgattccctc cagacctacg 2200  
 cttttgaggg aacagggta ttagctggat ccctgagctc cttagaatca 2250  
 gcagtctctg atcaggatga aagctatgat taccttaatg agttgggacc 2300  
 tcgcttttaa agattagcat gcatgtttgg ttctgcagtg cagtcaaata 2350  
 attagggctt tttacatca aaatttttaa aagtgcata gtgtattcga 2400  
 acccaatggg agtcttaaag agttttgtgc cctggctcta tggcggggaa 2450  
 agccctagtc tatggagttt tctgatttcc ctggagtaaa tactccatgg 2500  
 ttattttaag ctacctacat gctgtcattg aacagagatg tggggagaaa 2550  
 tgtaacaat cagctcacag gcatcaatac aaccagattt gaagtaaaat 2600  
 aatgtaggaa gatattaaaa gtagatgaga ggacacaaga tgtagtcgat 2650  
 ccttatgcga ttatatcatt atttacttag gaaagagtaa aaataccaaa 2700  
 cgagaaaatt taaaggagca aaaatttgca agtcaaataa aaatgtacaa 2750  
 atcgagataa catttacatt tctatcatat tgacatgaaa attgaaaatg 2800  
 tatagtcaga gaaattttca tgaattattc catgaagtat tgtttccttt 2850  
 atttaaa 2857

<210> 264

<211> 772

<212> PRT

<213> Homo sapiens

<400> 264

Met	Asn	Cys	Tyr	Leu	Leu	Leu	Arg	Phe	Met	Leu	Gly	Ile	Pro	Leu
1				5					10					15
Leu	Trp	Pro	Cys	Leu	Gly	Ala	Thr	Glu	Asn	Ser	Gln	Thr	Lys	Lys
				20					25					30
Val	Lys	Gln	Pro	Val	Arg	Ser	His	Leu	Arg	Val	Lys	Arg	Gly	Trp
				35					40					45
Val	Trp	Asn	Gln	Phe	Phe	Val	Pro	Glu	Glu	Met	Asn	Thr	Thr	Ser
				50					55					60